****
Reviewer Comments:
<210> 9390
<211>
<212> mRNA
<213> Artificial Sequence

<220>
<223> Antisense to TGF-beta1

<400> 9390
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ACATTTTTTAAAAGTGTTTTGTAGAGATAGGGTCTCACTATGTTACCCAGGCTGGTCTCAAATGCCTGGATT
CAAGTATCCTCCCATCTCTGCCTCCCAAAAGTGCTAGGATTACAGGCGTGAGCACCCCGCCTGGCCTGAACT
ACTATCTTTTATTGTCTTCTTCACTATCCCCCACTAAAGCAGGTTCCTGGTGGGCAGGAACTCCTCCCTTAA
CCTCTCTGGGCTTGTTTCCTCAACCTTTAAAATGGGTGTTATCAGAGTCCCTGCCATCTCAGAGTGTTGCTA
TGGTGACTGAATGAGTTCATTAATGTAAGGCACTTCAACAGTGCCCAAGGTGCTCAATAAATAGATCTAACT
ACAGTAGTGTTCCCCACTGGTCCCCTGTGCCTTGATGCCGGGCAAAGGAATAGTGCAGACAGGCAGGAGGAG
GCAGAGAGGGAGAGAGAGGGAGTGGGAGTGGGGGAACGTCAGGGATGGAGACCCCAGGCAGGCGCCCAATGA
CACAGAGATCCGCAGTCCTCTCTCCATCTTTAATGGGGCCCCAGGTGGGCTTGGGGCACGGTGTCCTTAAAT
ACAGCCCCCATGGGCAAGGCAGCGGGGGCGGGGCGGGGTGGGGCCGGGCCTGCCGGGGCGGGGCGGGGCGGG
GCGGGACCTCAGCTGCACTTGCAGGAGCGCACGATCATGTTGGACAGCTGCTCCACCTTGGGCTTGCGGCCC
ACGTAGTACACGATGGGCAGCGGCTCCAGCGCCTGCGGCACGCAGCACGGCGCCGCCGAGGCGCCCGGGTTA
TGCTGGTTGTACAGGGCCAGGACCTTGCTGTACTGCGTGTCCAGGCTCCAAATGTAGGGGCAGGGCCCGAGG
CAGAAGTTGGCATGGTAGCCCTTGGGCTCGTGGATCCACTTCCAGCCGAGGTCCTTGCGGAAGTCAATGTAC
AGCTGCCGCACGCAGCAGTTCTTCTCCGTGGAGCTGAAGCAATAGTTGGTGTCCAGGGCTCGGCGGTGCCGG
GAGCTTTGCAGATGCTGGGCCCTCTCCAGCGGGGTGGCCATGAGAAGCAGGAAAGGCCGGTTCATGCCATGA
ATGGTGGCCAGGTCACCTCGGCGGCCGGTAGTGAACCCGTTGATGTCCACTTGCAGTGTGTTATCCCTGCTG
TCACAGGAGCAGTGGGCGCTAAGGCGAAAGCCCTCAATTTCCCCTCCACGGCTCAACCACTGCCGCACAACT
CCGGTGACATCAAAAGATAACCACTCTGGCGAGTCGCTGGGTGCCAGCAGCCGGTTGCTGAGGTATCGCCAG
GAATTGTTGCTGTATTTCTGGTACAGCTCCACGTGCTGCTCCACTTTTAACTTGAGCCTCCTCAGCAGACGC

AGCTCTGCCCGGGAGAGCAACACGGGTTCAGGTACCGCTTCTCGGAGCTCTGATGTGTTGAAGAACATATAT
ATGCTGTGTGTACTCTGCTTGAACTTGTCATAGATTTCGTTGTGGGTTTCCACCATTAGCACGCGGGTGACC
TCCTTGGCGTAGTAGTCGGCCTCAGGCTCGGGCTCCGGTTCTGCACTCTCCCCGGCCACCGGTCGCGGGTG
CTGTTGTACAGGGCGAGCACGGCCTCGGGCAGCGGGCCGGGCGGCACCTCCCCCTGGCTCGGGGGGCTGGCG
AGCCGCAGCTTGGACAGGATCTGGCCGCGGATGGCCTCGATGCGCTTCCGCTTCACCAGCTCCATGTCGATA
GTCTTGCAGGTGGATAGTCCCGCGGCCGGCGGGCCAGGCGTCAGCACCAGTAGCCACAGCAGCGGTAGCAGC
AGCGGCAGCAGCCGCAGCCCGGAGGGCGGCATGGGGGAGGCGGCGCCCCCGGCACTGCCGAGAGCGCGAAC
AGGGCTGGTGTGGTGGGGAGGCCCCGCCCCTGCAGGGGCTGGGGGTCTCCCGGCAAAAGGTAGGAGGGCCTC
GAGGGAAAGCTGAGGCTCCTCAGGGAGAAGGGCGCAGTGGTGGAGGGGAGGCTTGGACCGGGGGTGTCTCAG
TATCCCACGGAAATAACCTAGATGGGCGCGATCTGGTACCAGAAGGTGGGTGGTCTTGAATAGGGGATCTGT
GGCAGGTCGGAGAGAGATCCGTCTCCTGGAGGAGAAAGGGTCTAGGATGCGCGGGGGCTCAGGAGACAGGCC
GGGGATGAAGGCGGCGTGCAGGGGGTGCGCCCGAGGTCTGGGGAAAAGTCTTTGCGGGAGGCCGGGTCGGCG
ACTCCCGAGGGCTGGTCCGGAATGGGGGCGCCTGAGGGACGCCGTGTAGGGGGCAGGGAGGGAGCAAGCGTC
CCCGGCGGCAAAGGGAGGCGGTCTGGGGTCCCCAAGTCCTGCCTCCTCGCGGGGCAGCGTCGCGCCAAGAGG
TCCCCGCGCCTCCGGCTCCCAGCGGCAACGGAAAAGTCTCAAAAGTTTTTTCCTCTTCTCCCGACCAGCTC
GTCCCTCCTCCCGCTCCTCCTCCCCCTCCTCCCCGCAGTGGCGGGGGCGGCGGCGGCTCGTCTCAGACTCTG
GGGCCTCAGGCTGCTCCTCGGCGACTCCTTCCTCCGCTCCGGGCCGAGGCCGGCCCCGCGGGCGGCTCAGAG
CCGGGGGGGGTGCCCCGGACGGGGCGTCCCCCCTGCCCCCGGCCGGGGCCCTCGCTGTCTGGCTGCTCCGCG
GAGGGAGGT

Mandatory field Numeric Identifier <211> is missing, which represents
the total number of bases in the sequence listing.

Nucleotide bases must be in groups of ten, Per Sequence Rules only 60
bases are allowed per line.

The sequence rules require that "the enumeration of nucleotide bases
shall start at the first base of the sequence with 1. The enumeration
shall be continuous through the whole sequence in the direction 5' to
3'. The enumeration shall be marked in the right margin, next to the
line containing the one-letter codes for bases, and giving the number of
the last base of that line."

Please change the nucleotides to lower-case letters. Do not use upper-
case letters for nucleotides. This error appears in subsequent
sequences in the sequence listing.

Please remove the two invalid carriage returns in the sequences.

Numeric Identifier <212> represent type of sequence DNA/RNA or PRT. If
the sequences are an RNA sequence, 't' are not allowed.

The above is a sample of multiple errors in the sequence listing, please make all necessary corrections.

* *

Application No: 10591048   Version No: 2.0

Input Set:

Output Set:

Started:	2010-05-18 14:49:04.181
Finished:	2010-05-18 15:08:14.478
Elapsed:	0 hr(s) 19 min(s) 10 sec(s) 297 ms
Total Warnings:	20192
Total Errors:	319
No. of SeqIDs Defined:	9394
Actual SeqID Count:	9394

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

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            Input Set:


            Output Set:



            Started:    2010-05-18 14:49:04.181
           Finished:    2010-05-18 15:08:14.478
            Elapsed:    0 hr(s) 19 min(s) 10 sec(s) 297 ms
     Total Warnings:    20192
       Total Errors:    319
 No. of SeqIDs Defined: 9394
   Actual SeqID Count:  9394
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Error code	Error Description
	This error has occured more than 20 times, will not be displayed
E 310	Invalid sequence type in <212> in SEQID: (9081)
E 310	Invalid sequence type in <212> in SEQID: (9082)
E 310	Invalid sequence type in <212> in SEQID: (9083)
E 310	Invalid sequence type in <212> in SEQID: (9084)
E 310	Invalid sequence type in <212> in SEQID: (9085)
E 310	Invalid sequence type in <212> in SEQID: (9086)
E 310	Invalid sequence type in <212> in SEQID: (9087)
E 310	Invalid sequence type in <212> in SEQID: (9088)
E 310	Invalid sequence type in <212> in SEQID: (9089)
E 310	Invalid sequence type in <212> in SEQID: (9090)
E 310	Invalid sequence type in <212> in SEQID: (9091)
E 310	Invalid sequence type in <212> in SEQID: (9092)
E 310	Invalid sequence type in <212> in SEQID: (9093)
E 310	Invalid sequence type in <212> in SEQID: (9094)
E 310	Invalid sequence type in <212> in SEQID: (9095)
E 310	Invalid sequence type in <212> in SEQID: (9096)
E 310	Invalid sequence type in <212> in SEQID: (9097)
E 310	Invalid sequence type in <212> in SEQID: (9098)
E 310	Invalid sequence type in <212> in SEQID: (9099)
E 310	Invalid sequence type in <212> in SEQID: (9100)
	This error has occured more than 20 times, will not be displayed
E 201	Mandatory field data missing in <211> in SEQ ID (9390)

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              Input Set:


              Output Set:



              Started:    2010-05-18 14:49:04.181
              Finished:   2010-05-18 15:08:14.478
              Elapsed:    0 hr(s) 19 min(s) 10 sec(s) 297 ms
      Total Warnings:     20192
        Total Errors:     319
No. of SeqIDs Defined:    9394
    Actual SeqID Count:   9394


Error code        Error Description


W  112        Upper case found in data; Found at position(0) SEQID(9390)

W  112        Upper case found in data; Found at position(1) SEQID(9390)

W  112        Upper case found in data; Found at position(2) SEQID(9390)

W  112        Upper case found in data; Found at position(3) SEQID(9390)

W  112        Upper case found in data; Found at position(4) SEQID(9390)

W  112        Upper case found in data; Found at position(5) SEQID(9390)

W  112        Upper case found in data; Found at position(6) SEQID(9390)

W  112        Upper case found in data; Found at position(7) SEQID(9390)

W  112        Upper case found in data; Found at position(8) SEQID(9390)

W  112        Upper case found in data; Found at position(9) SEQID(9390)

W  112        Upper case found in data; Found at position(10) SEQID(9390)

W  112        Upper case found in data; Found at position(11) SEQID(9390)

W  112        Upper case found in data; Found at position(12) SEQID(9390)

W  112        Upper case found in data; Found at position(13) SEQID(9390)

W  112        Upper case found in data; Found at position(14) SEQID(9390)

W  112        Upper case found in data; Found at position(15) SEQID(9390)

W  112        Upper case found in data; Found at position(16) SEQID(9390)

W  112        Upper case found in data; Found at position(17) SEQID(9390)

W  112        Upper case found in data; Found at position(18) SEQID(9390)

W  112        Upper case found in data; Found at position(19) SEQID(9390)
              This error has occured more than 20 times, will not be displayed

E  201        Mandatory field data missing in <211>  in SEQ ID (9391)

E  201        Mandatory field data missing in <211>  in SEQ ID (9392)
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        Input Set:


        Output Set:



          Started:   2010-05-18 14:49:04.181
         Finished:   2010-05-18 15:08:14.478
          Elapsed:   0 hr(s) 19 min(s) 10 sec(s) 297 ms
   Total Warnings:   20192
     Total Errors:   319
No. of SeqIDs Defined:   9394
 Actual SeqID Count:   9394
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Error code	Error Description
E 201	Mandatory field data missing in <211> in SEQ ID (9393)
E 201	Mandatory field data missing in <211> in SEQ ID (9394)

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<210> 2

<211> 14

<212> DNA

<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

<400> 2
gtcgatagtc ttgc   14

<210> 3

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 3
cttggacagg atct   14

<210> 4

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 4
ccaggaattg ttgc   14

<210> 5

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 5
cctcaatttc ccct   14

<210> 6

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 6
gatgtccact tgca   14

<210> 7

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

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ctccaaatgt aggg   14

<210> 8

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 8
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<210> 9

<211> 14

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gtagtacacg atgg   14

<210> 10

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<212> DNA

<213> Artificial Sequence

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<210> 11

<211> 14

<212> DNA

<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

<400> 11

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<210> 12

<211> 14

<212> DNA

<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

<400> 12

gcacgatcat gttg   14

<210> 13

<211> 16

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

<400> 13

tgtactctgc ttgaac   16

<210> 14

<211> 18

<212> DNA

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<223> Antisense to TGF-beta 1

<400> 14

ctgatgtgtt gaagaaca   18

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<211> 16

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<210> 16

<211> 16

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<210> 17

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

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<210> 18

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 1

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<210> 19

<211> 18

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<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

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gtcatagatt tcgttgtg   18

<210> 20

<211> 18

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<213> Artificial Sequence

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<223> Antisense to TGF-beta 1

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<210> 21

<211> 18

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<212>  DNA

<213>  Artificial Sequence



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<223>  Antisense to TGF-beta 2

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<210>  23

<211>  14

<212>  DNA

<213>  Artificial Sequence



<220>

<223>  Antisense to TGF-beta 2

<400>  23
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<210>  24

<211>  17

<212>  DNA

<213>  Artificial Sequence



<220>

<223>  Antisense to TGF-beta 2

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<210>  25

<211>  14
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<212> DNA

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<210> 26

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<212> DNA

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<223> Antisense to TGF-beta 2

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<210> 27

<211> 14

<212> DNA

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<223> Antisense to TGF-beta 2

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<210>   29

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<223>   Antisense to TGF-beta 2

<400>   29
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<210>   30

<211>   18

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<220>

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<210>   31

<211>   14

<212>   DNA

<213>   Artificial Sequence

<220>

<223>   Antisense to TGF-beta 2

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<210>   32

<211>   14

<212>   DNA

<213>   Artificial Sequence

<220>

<223>   Antisense to TGF-beta 2

<400>   32
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<210>   33

<211>   18

<212>   DNA

<213>   Artificial Sequence

<220>

<223>   Antisense to TGF-beta 2

<400>   33
acaattctga agtagggt   18

<210>   34

<211>   18

<212>   DNA

<213>   Artificial Sequence

<220>

<223>   Antisense to TGF-beta 2

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<210>   35

<211>   20

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<223> Antisense to TGF-beta 2

<400> 35
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<210> 36

<211> 14

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<220>

<223> Antisense to TGF-beta 2

<400> 36
ctggcttttg ggtt   14

<210> 37

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense to TGF-beta 2

<400> 37
tctgatatag ctcaatcc   18

<210> 38

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Antisense to TGF-beta 2

<400> 38
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<210> 39

<211> 16

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<223> Antisense to TGF-beta 2

<400> 39
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<210> 40

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<223> Antisense to TGF-beta 2

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<223> Antisense to TGF-beta 2

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<210> 42

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gcagcaatta tcctgc   16

<210> 43

<211> 14

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<223> Antisense to TGF-beta 2

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<210> 44

<211> 14

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<223> Antisense to TGF-beta 2

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<211> 18

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<223> Antisense to TGF-beta 2

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<210> 49

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<210> 50

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<220>

<223> Antisense to IL-10

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<210> 52

<211> 15

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<223> Antisense to IL-10

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<223> Antisense to IL-10

<400> 55
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<210> 56

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<223> Antisense to IL-10

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<223> Antisense to IL-10

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<210> 58

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<223> Antisense to IL-10

<400> 58
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<223> Antisense to IL-10

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<210> 60

<211> 18

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<210> 64

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<223> Antisense to TGF-beta 3

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<210> 72

<211> 14

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